FILE 'AGRICOLA, MEDLINE, CAPLUS, BIOSIS' ENTERED AT 09:37:32 ON 06 JUN

	2006		
L1		2185	S PENICILLIUM (2N) CITRINUM
L2		76	S L1 AND REDUCTASE
L3		21	S L1 (10N) REDUCTASE
L4		16	DUP REM L3 (5 DUPLICATES REMOVED)
L5		9	S L4 AND PY<2003
L6		4	S ASAKO
L7		1006	S ASAKO OR SHIMIZU
L8		0	S L7 AND CITRINUM

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L5
     ANSWER 1 OF 9
                      MEDLINE on STN
                MEDLINE
AN
     82007510
DN
     PubMed ID: 7275844
ΤI
     Dihydrocompactin, a new potent inhibitor of 3-hydroxy-3-methylglutaryl
     coenzyme-A reductase from Penicillium citrinum
ΑU
     Lam Y K; Gullo V P; Goegelman R T; Jorn D; Huang L; DeRiso C; Monaghan R
     L; Putter I
SO
     The Journal of antibiotics, (1981 May) Vol. 34, No. 5, pp.
     614-6.
     Journal code: 0151115. ISSN: 0021-8820.
CY
     Japan
DT
     Journal; Article; (JOURNAL ARTICLE)
LA
     English
FS
     Priority Journals
EM
     198111
ED
     Entered STN: 16 Mar 1990
     Last Updated on STN: 16 Mar 1990
     Entered Medline: 18 Nov 1981
     ANSWER 2 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN
AN
     2002:446160 CAPLUS
DN
     137:19459
TI
    Manufacture of optically active 4-halo-3-hydroxybutanoate by asymmetric
     reduction of 4-halo-3-oxobutanoate with transgenic microorganisms
     Asako, Hiroyuki; Matsumura, Kenji; Shimizu, Masatoshi; Ito, Nobuya;
IN
     Wakita, Ryuhei
PA
     Sumitomo Chemical Company, Limited, Japan
SO
     Eur. Pat. Appl., 56 pp.
     CODEN: EPXXDW
DT
     Patent
     English
LΑ
FAN. CNT 1
     PATENT NO.
                    KIND
                               DATE APPLICATION NO. DATE
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                               -----
                                           -----
                                                                  _____
    EP 1213354
                        A2
PΙ
                               20020612 EP 2001-310251 20011207 <--
     EP 1213354
                        A3
                               20030205
         R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
             IE, SI, LT, LV, FI, RO, MK, CY, AL, TR
    JP 2002212158 A2
JP 200223789 A2
JP 2002233392 A2
JP 2003061682 A2
US 2003134402 A1
                               20020731
                                          JP 2001-6144
                                                                  20010115 <--
                               20020813
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                                                                  20010202 <--
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     US 6884607
                       B2
                               20050426
US 6884607
US 2005019816 A1
PRAI JP 2000-372704 A
                                          US 2004-920239
                               20050127
                                                                 20040818
                               20001207
     JP 2001-6144
                        Α
                               20010115
     JP 2001-26594
                         Α
                               20010202
     JP 2001-175175
                         Α
                               20010611
     US 2001-4115
                         A3
                               20011206
os
     CASREACT 137:19459; MARPAT 137:19459
AB
     A method of manufacturing of optically active 4-halo-3-hydroxybutanoic acids
     from the corresponding 4-halo-3-oxobutanoate using by stereospecific
     enzymic reduction is described. The method uses a reductase from
     Penicillium citrinum. The gene for the enzyme is
     expressed in a transgenic host such as Escherichia coli. The enzyme was
     purified chromatog. from mycelium of Penicillium citrinum and amino acid
     sequence-derived primers used to clone the gene. Escherichia coli
     expressing the gene for the enzyme was used to reduce 4-bromo-3-
     oxobutanoate to methyl-(S)-4-Bromo-3-hydroxybutanoate. Cells, 0.4 g wet
     weight, were incubated with Me 4-bromo-3-oxobutanoate 1500 mg in the presence
     of NADP+ 18 mg, glucose 3000 mg, glucose dehydrogenase 3 mg, in 15 mL of
```

phosphate buffer pH 6.5 and Bu acetate 15 mL at 30° for 7 h. Yield was 99.2% with an ee of 95.7%.

- L5 ANSWER 3 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN
- AN 2001:137378 CAPLUS
- DN 134:173918
- TI Penicillium citrinum genes associated with biosynthesis of ML-236B, precursor of a 3-hydroxy-3-methylglutaryl CoA reductase inhibitor
- IN Abe, Yuki; Hosobuchi, Masahiko; Yoshikawa, Hiroji
- PA Sankyo Company, Ltd., Japan
- SO PCT Int. Appl., 128 pp.

CODEN: PIXXD2

DT Patent

LA Japanese

FAN.CNT 1

PATENT NO. KIND DATE APPLICATION NO. DATE ---------_____ -----20010222 WO 2000-JP5420 PΙ WO 2001012814 A1 20000811 <--W: AU, BR, CA, CN, CZ, HU, ID, IL, IN, KR, MX, NO, NZ, PL, RU, TR, US, ZA RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE

JP 2001112487 A2 20010424 PRAI JP 1999-227696 A 19990811

AB Genes associated with biosynthesis of ML-236B, a precursor of pravastatin, a 3-hydroxy-3-methylglutaryl CoA (HMG-CoA) reductase inhibitor, from Penicillium citrinum, are disclosed. PCR primers corresponding to those gene sequences are claimed. A gene cluster containing 6 genes, named mlcA, mlcB, mlcC, mlcD, mlcE, and mlcR, was cloned from Penicillium citrinum SANK13380 strain and sequenced. Penicillium citrinum strains transformed with those genes and having improved ML-236B production (average 12%) were obtained.

JP 2000-240722

20000809 <--

- RE.CNT 14 THERE ARE 14 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L5 ANSWER 4 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN
- AN 1994:75515 CAPLUS
- DN 120:75515
- TI Production of ML-236B, and inhibitor of 3-hydroxy-3-methylglutaryl CoA reductase by Penicillium citrinum: improvements of strain and culture conditions
- AU Hosobuchi, Masahiko; Shioiri, Tetsuya; Ohyama, Jo; Arai, Masatoshi; Iwado, Seigo; Yoshikawa, Hiroji
- CS Ferment. Res. Lab., Sankyo Co., Ltd., Iwaki, 970-04, Japan
- SO Bioscience, Biotechnology, and Biochemistry (1993), 57(9), 1414-19
 - CODEN: BBBIEJ; ISSN: 0916-8451
- DT Journal
- LA English
- AB In order to increase the yield of ML-236B, an intermediate for pravastatin Na (an inhibitor of cholesterol synthesis) production, the improvements of an ML-236B-producing strain of P. citrinum, the medium composition, and the culture conditions were studied. A mutant strain S-5808, which produces 20-fold more ML-236B than the original strain does in flask culture, was isolated. As S-5808 required a large amount of the C source for production of ML-236B, a fed-batch culture method was developed. The continuous feeding of the C source (glycerol or mixture of glycerol and maltose) was very effective for high ML-236B production ML-236B in the culture broth became oily with the decreasing pH during cultivation, adhered to the cells, decreased substrate consumption, and finally stopped ML-236B production late in fermentation The addition of a polypropylene glycol-type surfactant enhanced

ML-236B productivity.

L5 ANSWER 5 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN

AN 1992:248436 CAPLUS

DN 116:248436

TI Aldose reductase inhibitors contain citrinin derivatives or dihydrocitrinin derivatives for treatment of diabetic complication

IN Takemura, Ayako; Yamamoto, Hideki; Donho, Munehiko

PA Unitika Ltd., Japan

SO Jpn. Kokai Tokkyo Koho, 5 pp.

CODEN: JKXXAF

DT Patent

LA Japanese

FAN.CNT 1

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
ΡI	JP 04009332	A2	19920114	JP 1990-111259	19900425 <
PRAI	JP 1990-111259		19900425		
GI					

AB Aldose reductase inhibitors contain citrinin derivs. I or dihydrocitrinin derivs. II [R = H, R1, COR1 (R1 = alkyl, alkenyl, aryl, aralkyl); X = OR2, NR2R3 (R2, R3 = H, alkyl, alkenyl, aryl, aralkyl)], which are useful for prevention and treatment of diabetic complication. Penicillium citrinum (IFO 4631) was aerobically cultured in a medium containing glucose, meat extract,

peptone, and NaCl at 28° for 5 days to produce citrinin (III). IC50 of III against aldose reductase was 3.4 + 10-6M.. III at 200 mg/kg p.o. did not cause death in mice.

L5 ANSWER 6 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN

AN 1987:192494 CAPLUS

DN 106:192494

TI Effect of nitrate reductase on the reduction of iron

AU Shakhobova, B. B.

CS Nauchno-Issled. Inst. Pochvoved., Dushanbe, USSR

SO Izvestiya Akademii Nauk Tadzhikskoi SSR, Otdelenie Biologicheskikh Nauk (
1986), (2), 84-6

CODEN: ITOBAO; ISSN: 0002-3477

DT Journal

LA Russian

AB Eight strains of aerobic bacteria and fungi were studied for their ability to reduce NO3- and Fe(OH)3. There was no correlation between the 2 parameters. Asperigillus niger, Fusarium oxysporum And 2 strains of Arthrobacter aurescens reduced Fe(OH)3 but had no effect on NO3-. In contrast, Streptomyces albus reduced NO3- but had no effect on Fe3+. Penicillium glaucum And P. citrinum showed variable reactions. One strain of A. aurescens reduced Fe3+ but not NO3-.

L5 ANSWER 7 OF 9 CAPLUS COPYRIGHT 2006 ACS on STN

AN 1977:85285 CAPLUS

DN 86:85285

TI Competitive inhibition of 3-hydroxy-3-methylglutaryl coenzyme A reductase

by ML-236A and ML-236B fungal metabolites, having hypocholesterolemic activity

- AU Endo, Akira; Kuroda, Masao; Tanzawa, Kazuhiko
- CS Fermentation Res. Lab., Sankyo Co., Ltd., Tokyo, Japan
- SO FEBS Letters (1976), 72(2), 323-6 CODEN: FEBLAL; ISSN: 0014-5793
- DT Journal
- LA English

GI

HO2CCH2CH (OH) CH2CH (OH) CH2CH2 OR

AB Fungal metabolites, ML-236A (I) and ML-236B (II), from Penicillium citrinum inhibited 3-hydroxy-3-methylglutaryl-CoA reductase, the rate-limiting enzyme in the pathway of cholesterol synthesis. The acid forms (Na salts) of both compds. were more potent inhibitors than the lactone forms and both forms of II were more effective than those of I. Inhibition was competitive with respect to 3-hydroxy-3-methylglutaryl-CoA and noncompetitive with respect to NADPH. The Ki values were 0.22 μM for I Na salt and 0.010 μM for II Na salt.

- L5 ANSWER 8 OF 9 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
- AN 1982:103371 BIOSIS
- DN PREV198223033363; BR23:33363
- TI TOTAL SYNTHESIS OF DEXTRO COMPACTIN ML-236B.
- AU WANG N-Y [Reprint author]; HSU C-T; SIH C J
- CS SCH PHARM, UNIV WIS, MADISON, WIS 53706, USA
- 50 Journal of the American Chemical Society, (1981) Vol. 103, No. 21, pp. 6538-6539.
 CODEN: JACSAT. ISSN: 0002-7863.
- DT Article
- FS BR
- LA ENGLISH
- L5 ANSWER 9 OF 9 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
- AN 1979:266972 BIOSIS
- DN PREV197968069476; BA68:69476
- TI HYPO LIPIDEMIC EFFECTS IN MONKEYS OF ML-236B A COMPARATIVE INHIBITOR OF 3 HYDROXY-3-METHYL GLUTARYL COENZYME A REDUCTASE.
- AU KURODA M [Reprint author]; TSUJITA Y; TANZAWA K; ENDO A
- CS FERMENT RES LAB, SANKYO CO LTD, 1-2-58 HIROMACHI, SHINAGAWA, TOKYO 140, JPN
- SO Lipids, (1979) Vol. 14, No. 6, pp. 585-589. CODEN: LPDSAP. ISSN: 0024-4201.
- DT Article
- FS BA
- LA ENGLISH
- AB The fungal metabolite ML-236B from Penicillium citrinum, a competitive inhibitor of 3-hydroxy-3-methylglutaryl CoA reductase, was effective in lowering serum cholesterol levels in cynomolgus monkeys at doses of 20-50 mg/kg per day. Levels of serum phospholipids and triglycerides were not significantly changed by the administration of the drug. Of the serum lipoprotein fractions a β-lipoprotein corresponding to low density lipoprotein was preferentially reduced by the drug treatment. Fecal excretion of neutral

sterols was unaffected, but that of bile acids was slightly elevated by the administration of ML-236B.

Inventor Search

	ט	1	Document ID	Kind Codes	Source	Issue Date	Pages
1	х		US 20050191738 A1	IVII AD K LA LOS	US- PGPUB	20050901	23
2	х		US 20050063994 A1	ו עבאוז מומו	US- PGPUB	20050324	117
3	х		US 20050019816 A1	10/920,239	US- PGPUB	20050127	41
4	х		US 20040091902 A1	instant	US- PGPUB	20040513	16
5	X		US 20030186400 A1		US- PGPUB	20031002	26
6	х		US 20030134402 A1	10/9/125	US- PGPUB	20030717	41
7	х		US 20030039660 A1	10/ 44/15	US- PGPUB	20030227	114
8	х		US 6884607 B2	×	USPAT	20050426	37
9	Х		US 5264484 A	x	USPAT	19931123	5
10	Х		US 4229539 A	Х	USPAT	19801021	6
11	x		US 4127654 A	*	USPAT	19781128	13
12	х		JP 2002223789 A		JPO	20020813	6
13	х		JP 11164683 A		JPO	19990622	5
14	Х		JP 04365499 A		JPO	19921217	7
15	х		EP 1378566 A2	A2, A3	EPO	20040107	28
16	х		EP 1213354 A	245 or 271	DERWENT	20020612	37

	Title
1	Modified reductase and its gene, and use thereof
2	Methods and reagents for decreasing clinical reaction to allergy
3	Process for producing optically active-4-halo- 3-hydroxybutanoate
4	Modified reductase and its gene
5	Method for producing optically active 2- hydroxycycloalkanecarboxylic acid ester
6	Process for producing optically active 4-halo- 3-hydroxybutanoate
7	Recombinant hybrid allergen constructs with reduced allergenicity that retain immunogenicity of the natural allergen
8	Process for producing optically active 4-halo- 3-hydroxybutanoate
9	Room temperature curable organopolysiloxane composition having mildewproofing property
10	.betaGalactosidase and production thereof
11	Compositions and methods containing beta substituted allyl alcohols, sulfuric acid esters thereof, phosphoric acid esters thereof, alkanoyl esters thereof and alkylene oxide ethers thereof
12	METHOD FOR PRODUCING OPTICALLY ACTIVE 3- HYDROXYBUTYRIC ESTER COMPOUND
13	PROMOTION OF INFECTION OF PLANT BELONGING TO GENUS EUCALYPTLIS WITH MYCORRHIZAL FUNGUS AND INFECTION PROMOTER
14	INDICATOR FOR HIGH-PRESSURE STERILIZATION
15	Modified reductase from Penicillium citrinum and its encoding gene
16	New protein preferentially producing (S)-4- bromo-3-hydroxybutanoate by asymmetrically reducing 4-bromo-3-oxobutanoate, useful in pharmaceuticals and agrochemicals

	Туре	Hits	Search Text
1	BRS	1	US20030134402A1
2	BRS	2	"6884607"
3	BRS	474	penicillium near1 citrinum
4	BRS	139	S103 and reductase
5	BRS	2155	asako
6	BRS	10	S105 and citrinum
7	BRS	616	"110" and "245"
8	BRS	4	S106 and "245"
9	BRS	12	L5 near10 reductase
10	BRS	474	penicillium near1 citrinum
11	BRS	16	13 and citrinum
12	BRS	142777	shimizu or asako
13	BRS	14	11 and citrinum
14	BRS	140765	shimizu
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18	BRS	2	"20050272136"

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3	574.5	33.2	313	2	US-09-734-237B-44	Sequence 44, Appl
4	558.5	32.3	290	2	US-09-248-796A-17316	Sequence 17316, A
5	555	32.1	312	2	US-09-487-558B-438	Sequence 438, App
6	555	32.1	313	2	US-09-734-237B-39	Sequence 39, Appl
7	547.5	31.6	344	2	US-09-630-983A-20	Sequence 20, Appl
8	547	31.6	328	2	US-10-040-416-6	Sequence 6, Appli
9	545.5	31.5	328	2	US-10-040-416-4	Sequence 4, Appli
10	543.5	31.4	313	2	US-09-347-803-6	Sequence 6, Appli
11	541	31.3	322	2	US-09-734-237B-32	-

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 16, 2005, 19:49:35; Search time 189 Seconds

(without alignments)

755.546 Million cell updates/sec

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Searched: 2443163 seqs, 439378781 residues

Total number of hits satisfying chosen parameters: 2443163

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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3	1731	100.0	325	8	ADH44214	Adh44214 Penicilli
4	1731	100.0	325	8	ADL18258	Adl18258 Penicilli
5	1731	100.0	325	8	ADK70247	Adk70247 Penicilli
6	1731	100.0	325	8	ADM46686	Adm46686 325 amino
7	1731	100.0	325	8	ADM46567	Adm46567 Penicilli
8	1731	100.0	325	8	ADK51190	Adk51190 Wild-type
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10	1728	99.8	325	8	ADK51203	Adk51203 Mutant Pe
11	1726	99.7	325	8	ADK51204	Adk51204 Mutant Pe
12	1723	99.5	325	8	ADK51205	Adk51205 Mutant Pe
13	972.5	56.2	255	8	ADN20743	Adn20743 Bacterial

14	575.5	33.2	312	2	AAW29220	Aaw29220 S. cerevi
15	574.5	33.2	312	2	AAW29217	Aaw29217 S. cerevi
16	574.5	33.2	312	4	AAG63565	Aag63565 Amino aci
17	574.5	33.2	312	5	ABG93198	Abg93198 S. cerevi
18	574.5	33.2	312	8	ADS43942	Ads43942 Bacterial
19	574.5	33.2	313	4	AAG63566	Aag63566 Synthetic
20	573.5	33.1	312	2	AAW29218	Aaw29218 S.

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Published Applications AA Main:*
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2	1731	100.0	325	4	US-10-327-108-3	Sequence 3, Appli
3	1731	100.0	325	4	US-10-608-533-1	Sequence 1, Appli
4	1731	100.0	325	5	US-10-920-239-1	Sequence 1, Appli
5	1731	100.0	325	5	US-10-608-625-1	Sequence 1, Appli
6	994.5	57.5	309	4	US-10-425-115-287371	Sequence 287371,
7	972.5	56.2	255	4	US-10-369-493-3396	Sequence 3396, Ap
8	881	50.9	224	4	US-10-425-115-240890	Sequence 240890,
9	659	38.1	185	4	US-10-425-115-332876	Sequence 332876,
10	574.5	33.2	312	3	US-09-734-237B-42	Sequence 42, Appl
11	574.5	33.2	312	4	US-10-369-493-22372	Sequence 22372, A
12	574.5	33.2	312	4	US-10-451-467A-354	Sequence 354, App
13	574.5	33.2	312	5	US-10-989-488A-42	Sequence 42, Appl
14	574.5	33.2	313	3	US-09-734-237B-44	Sequence 44, Appl
15	574.5	33.2	313	5	US-10-989-488A-44	Sequence 44, Appl
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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-	1	1731	100.0	325	 6	US-10-617-034A-3	Seguence 3, Appli
	2	547.5	31.6	344	6	US-10-606-302-20	Sequence 20, Appl
	3	486.5	28.1	316	6	US-10-821-234-1526	Sequence 1526, Ap
	4	457	26.4	323	6	US-10-878-556A-13	Sequence 13, Appl
	5	454	26.2	323	6	US-10-878-556A-12	Sequence 12, Appl
	6	448	25.9	285	7	US-11-074-176-206	Sequence 206, App
	7	428.5	24.8	305	7	US-11-055-822-860	Sequence 860, App
	8	423.5	24.5	364	6	US-10-623-155-172	Sequence 172, App
	9	423.5	24.5	371	6	US-10-623-155-433	Sequence 433, App
	10	393	22.7	269	7	US-11-055-822-856	Sequence 856, App
	11	374.5	21.6	232	6	US-10-793-626-2194	Sequence 2194

PIR_80:* Database : 1: pir1:* 2: pir2:* 3: pir3:*

4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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2	568	32.8	321	2	T38413	probable oxidoredu
3	559.5	32.3	323	2	s78113	aldehyde reductase
4	555	32.1	312	2	S61163	aldo-keto reductas
5	553	31.9	327	1	s76143	probable aldehyde
6	548.5	31.7	325	2	T39169	probable oxidoredu
7	547.5	31.6	344	1	S46020	probable aldehyde
8	529.5	30.6	309	2	A84599	hypothetical prote
9	529	30.6	290	2	T02543	aldehyde dehydroge
10	527.5	30.5	309	2	B84599	hypothetical prote
11	523.5	30.2	294	2	T02542	probable alcohol d
12	523.5	30.2	313	2	T09670	abscisic

Database : UniProt_05.80:*

1: uniprot_sprot:* 2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		8			•	
Result		Query				
No.	Score	Match	Length	DB	ID	Description
1	1514	87.5	325	2	Q877A2_ASPOR	Q877a2 aspergillus
2	1477	85.3	325	2	Q4WQ65_ASPFU	Q4wq65 aspergillus
3	1471	85.0	325	2	Q5B1L7_EMENI	Q5b1l7 aspergillus
4	1471	85.0	325	2	Q7Z8L1_EMENI	Q7z8ll emericella
5	1205.5	69.6	331	2	Q7S3U4 NEUCR	Q7s3u4 neurospora
6	1191	68.8	327	2	Q6RZX1_TRIAT	Q6rzx1 trichoderma
7	1137.5	65.7	323	2	Q51QM9 MAGGR	Q51qm9 magnaporthe
8	1038	60.0	256	2	Q4I4F0_GIBZE	Q4i4f0 gibberella
9	1037	59.9	254	2	O74646_GIBZE	074646 gibberella
10	635	36.7	355	2	Q4PHK0_USTMA	Q4phk0 ustilago ma
11	623	36.0	1224	2	Q4P7C0 USTMA	Q4p7c0 ustilago ma
12	602.5	34.8	332	2	Q4IEY5_GIBZE	Q4iey5 gibberella
13	595.5	34.4	309	2	Q55SW0 CRYNE	Q55sw0 cryptococcu
14	593.5	34.3	309	2	Q5KH94 CRYNE	Q5kh94 cryptococcu
15	583	33.7	310	2	Q6FR42 CANGA	Q6fr42 candida gla
16	575.5	33.2	310	2	Q6FY54 CANGA	Q6fy54 candida gla
17	574.5	33.2	312	1	GCY YEAST	P14065 saccharomyc
18	570.5	33.0	309	2	Q6CRC8_KLULA	Q6crc8